



1/14

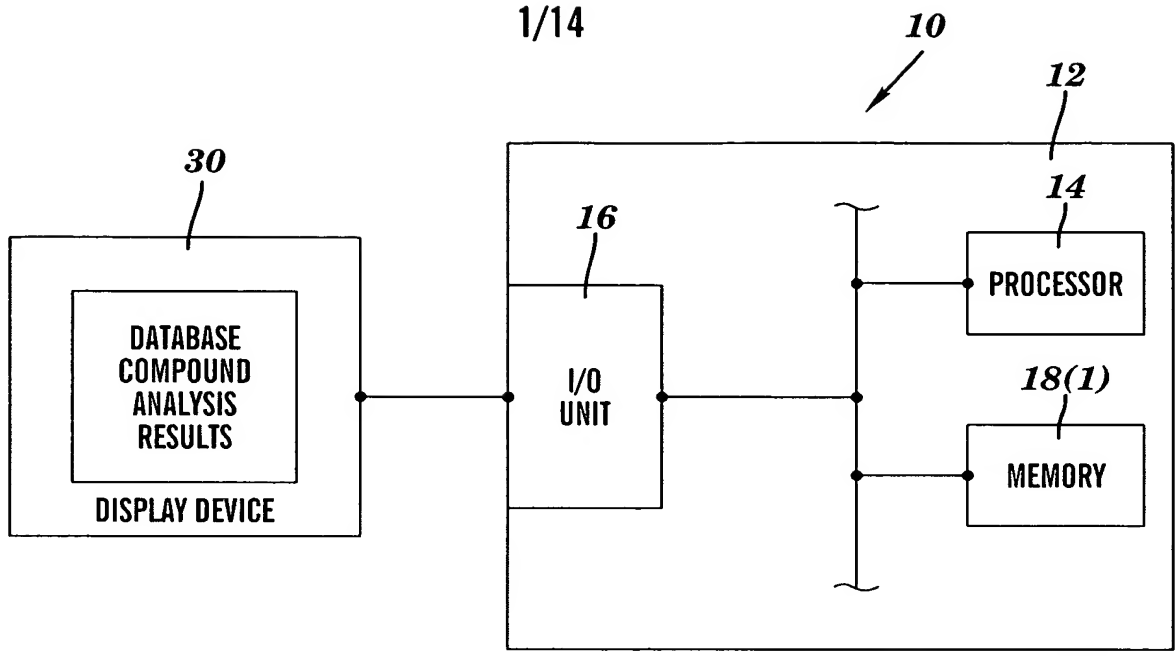


FIG. 1

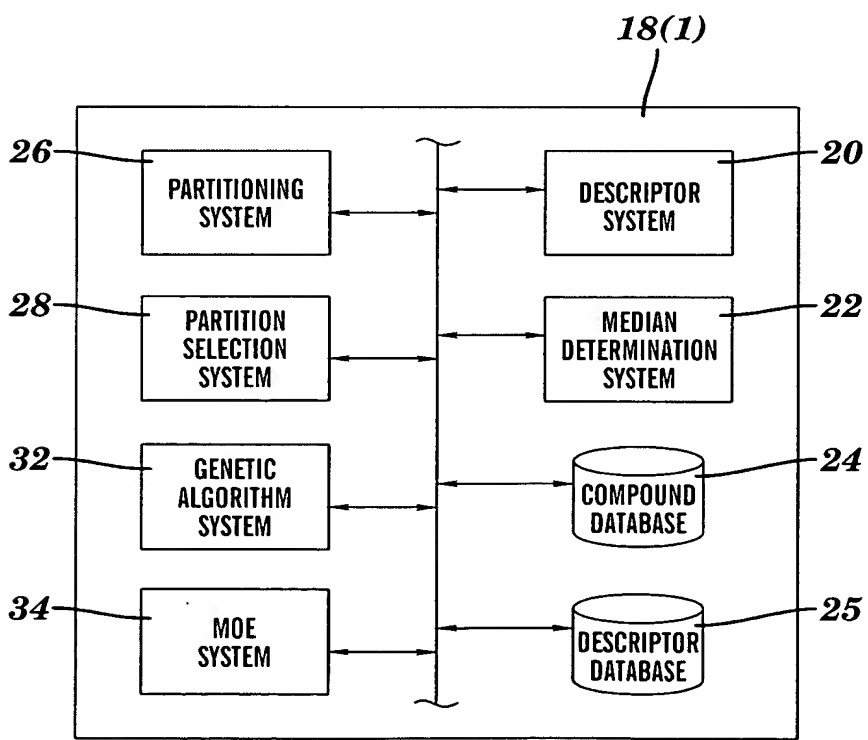
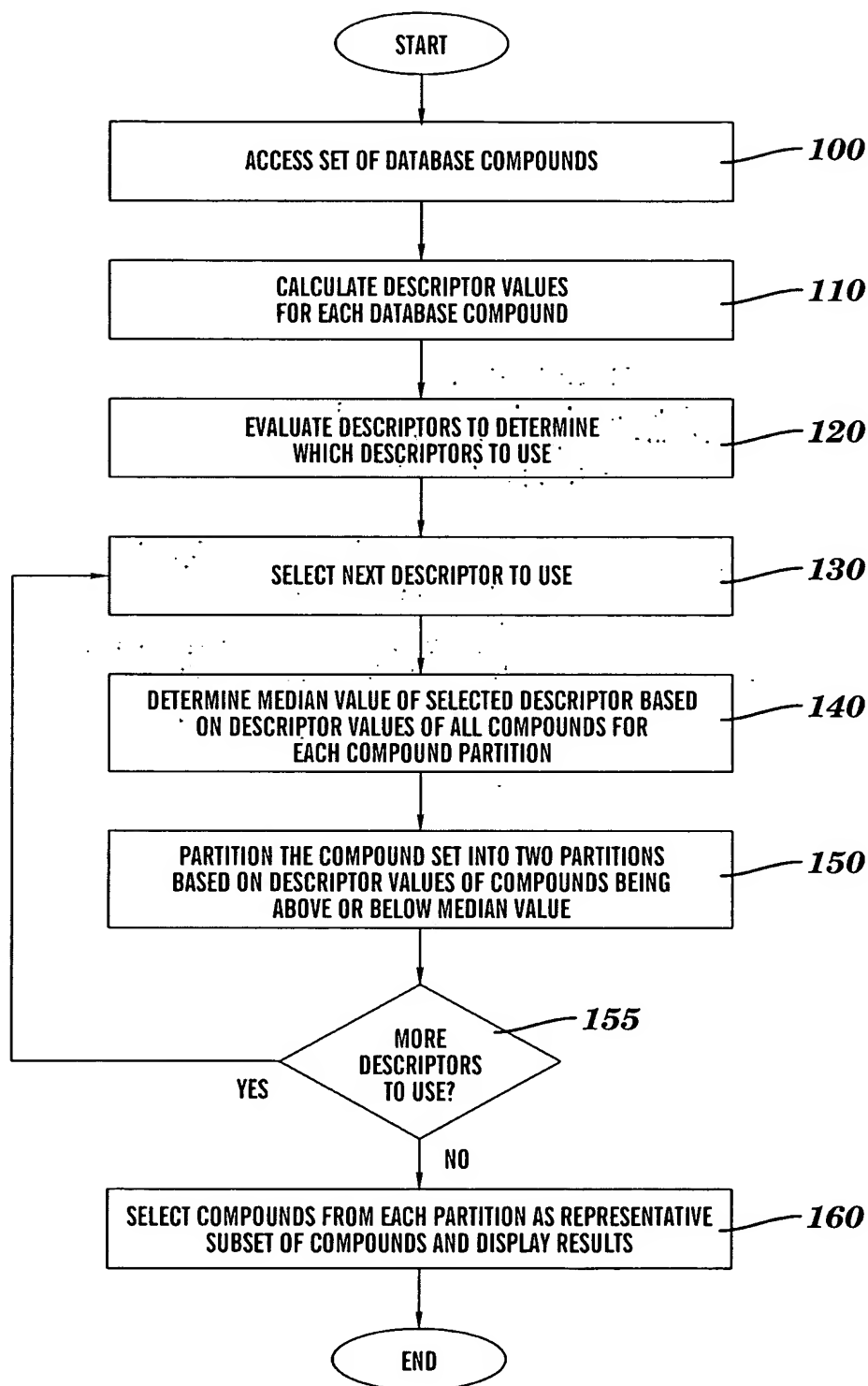


FIG. 2

**FIG. 3**

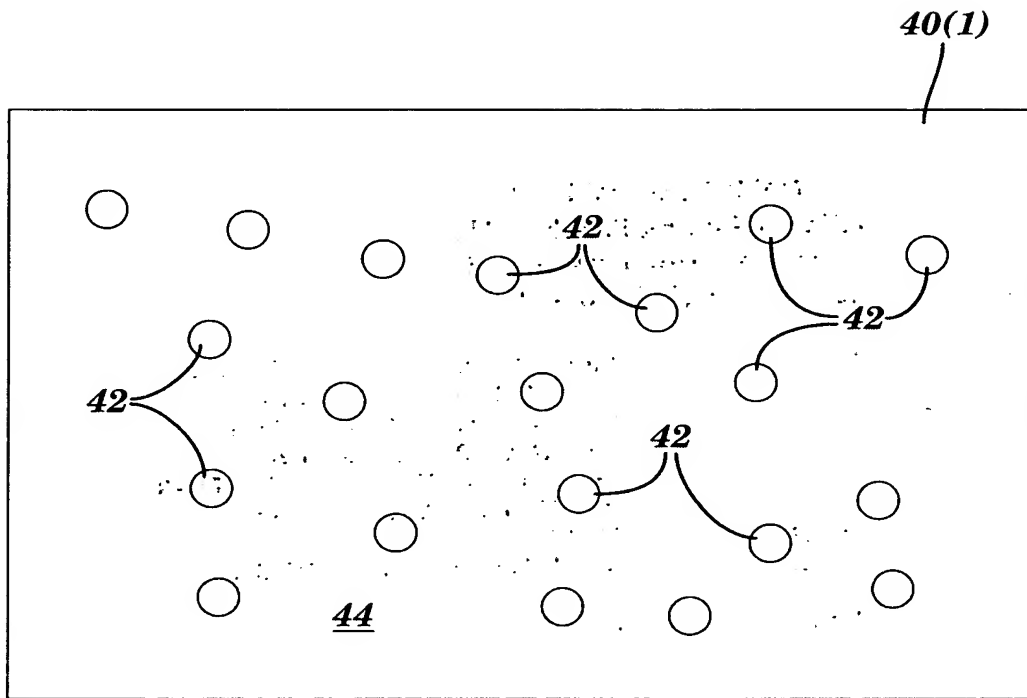
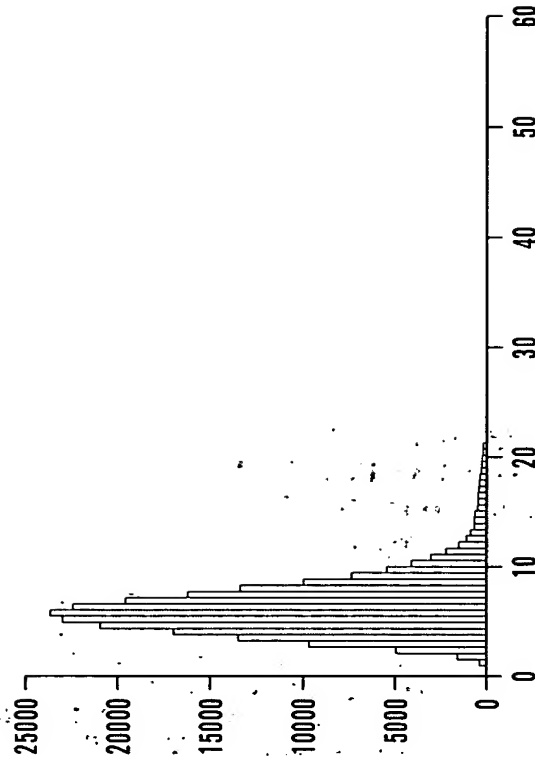


FIG. 4

MEAN: 6.17
MEDIAN: 5.81
SE: 0.62

KierA2



MEAN: 306
MEDIAN: 301
SE: 0.751

MW

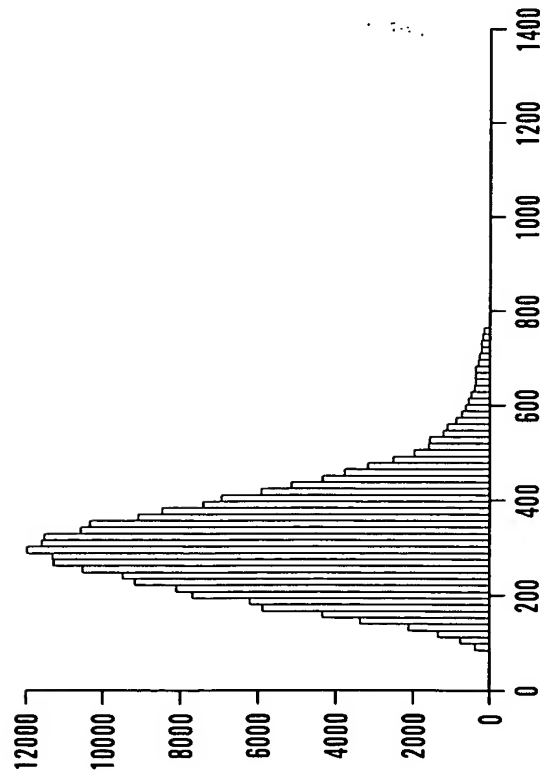
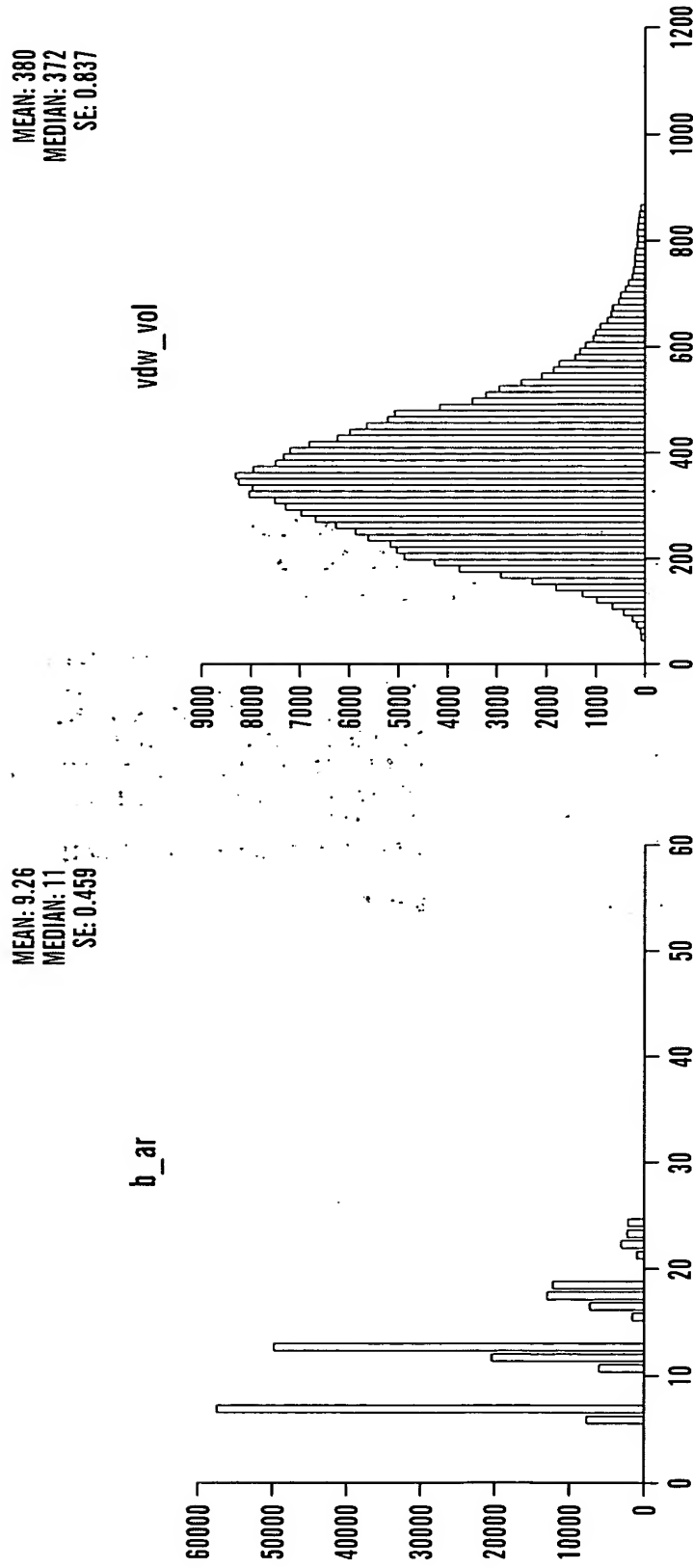


FIG. 5B

FIG. 5A



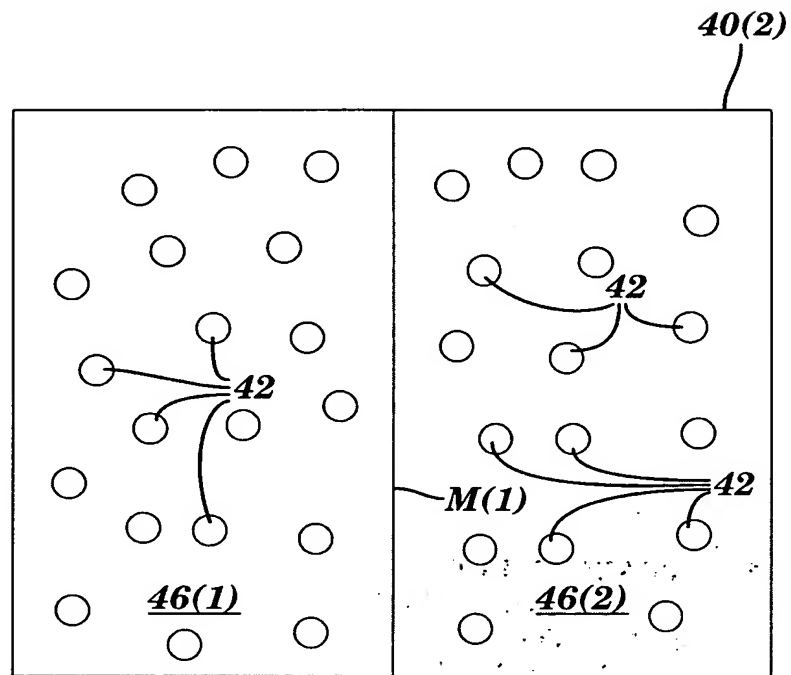


FIG. 6

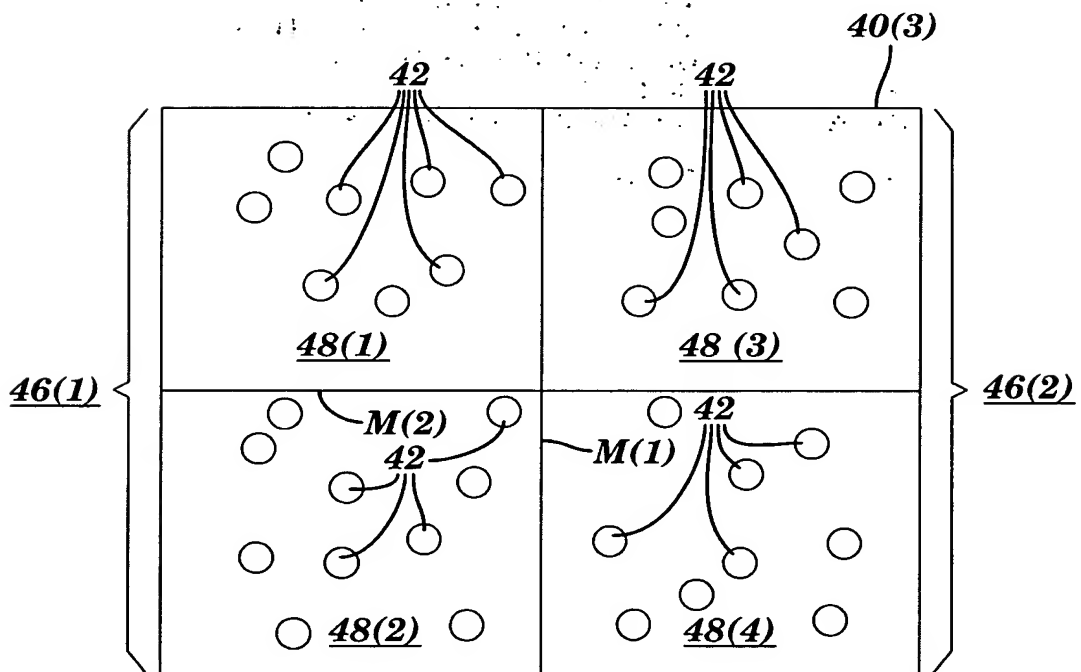


FIG. 7

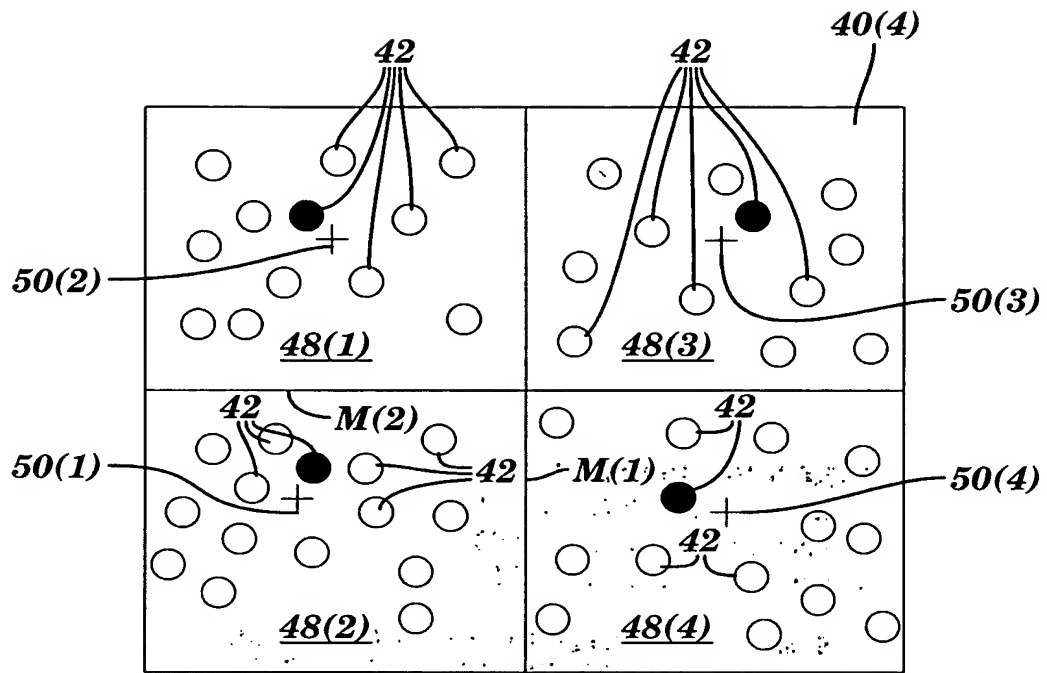
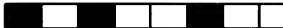




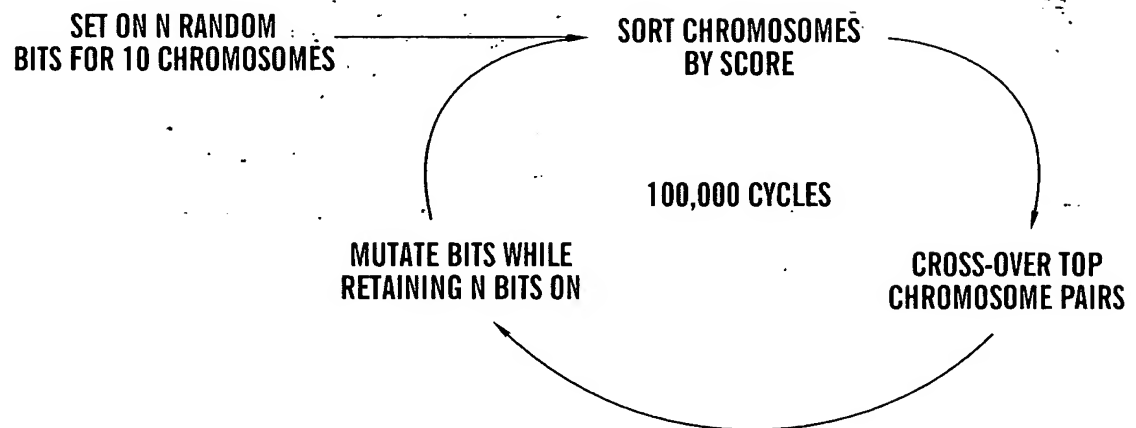


FIG. 8

CHROMOSOMES		AVERAGE SE	AVERAGE CORRELATION	SCORE
	• • •	0.476	0.131	3.634
	• • •	0.372	0.293	1.270
	• • •	0.611	0.535	1.342
	• • •	0.277	0.719	0.385
	• • •	0.198	0.886	0.223
← 111 DESCRIPTORS →				

**FIG. 9**

9/14

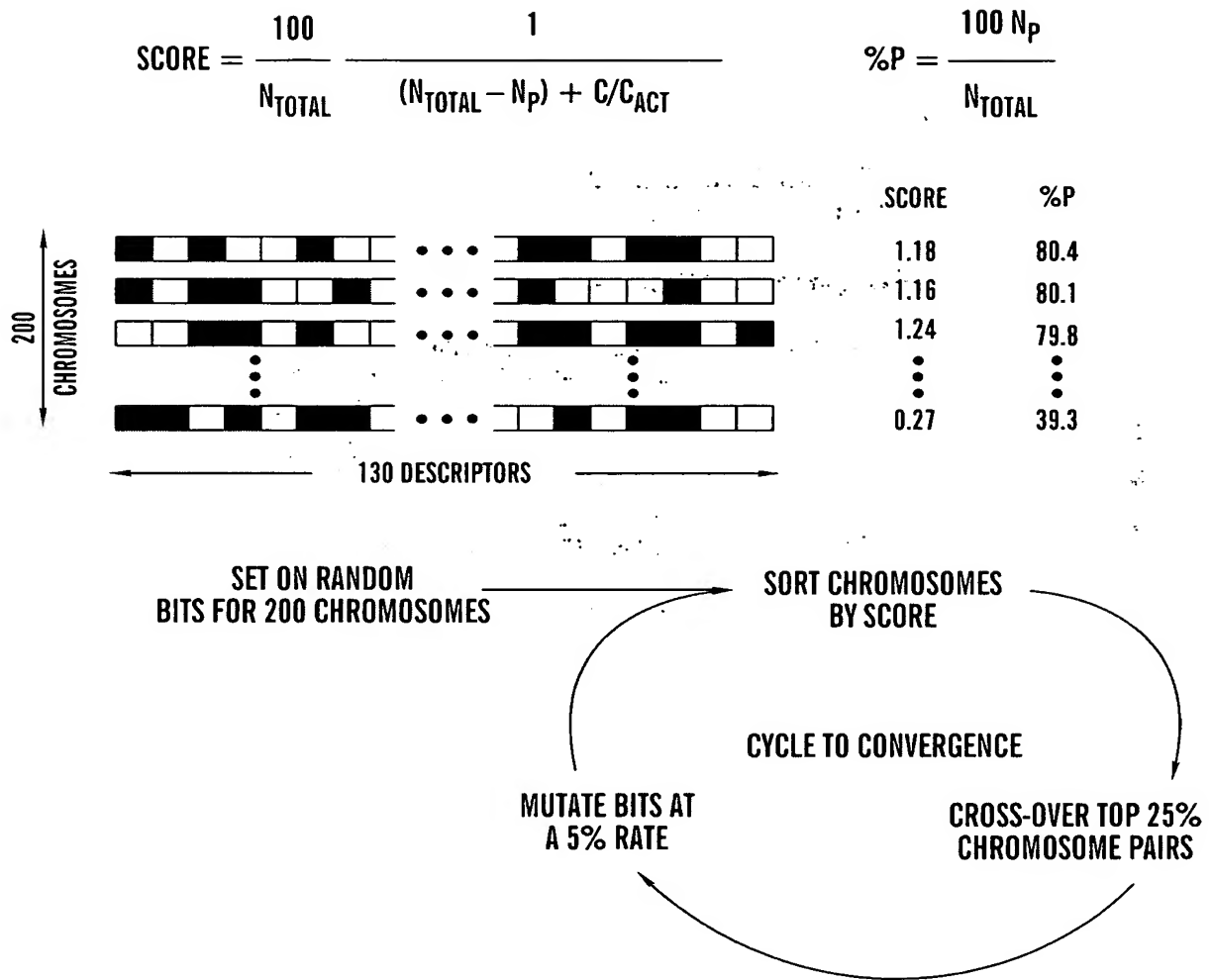
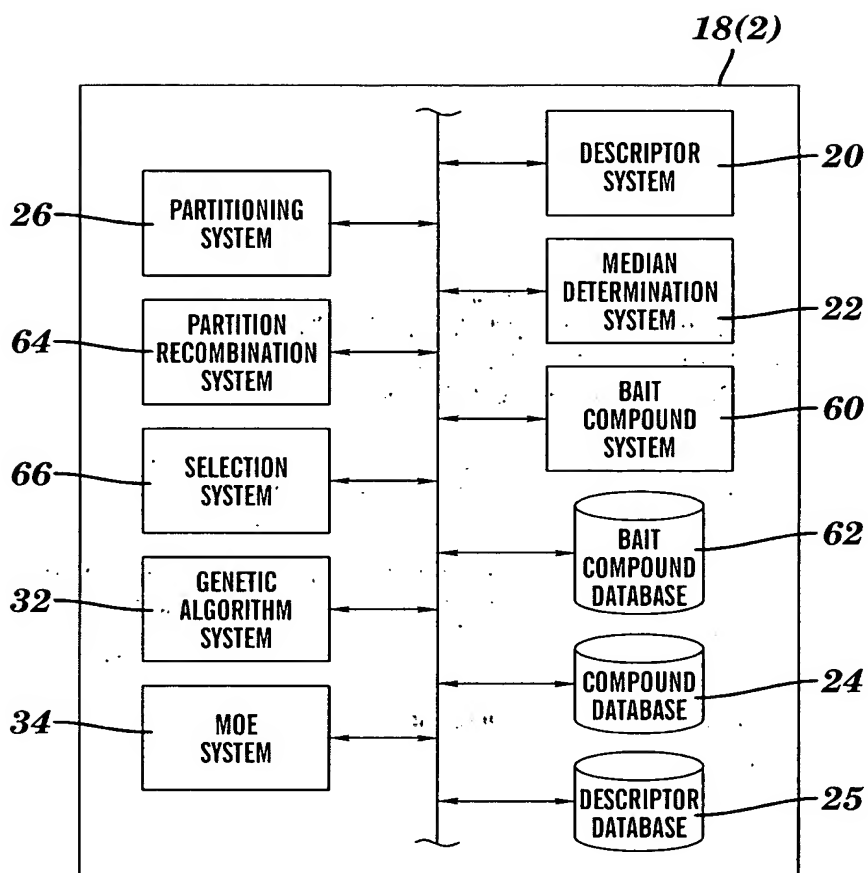


FIG. 10

**FIG. 11**

11/14

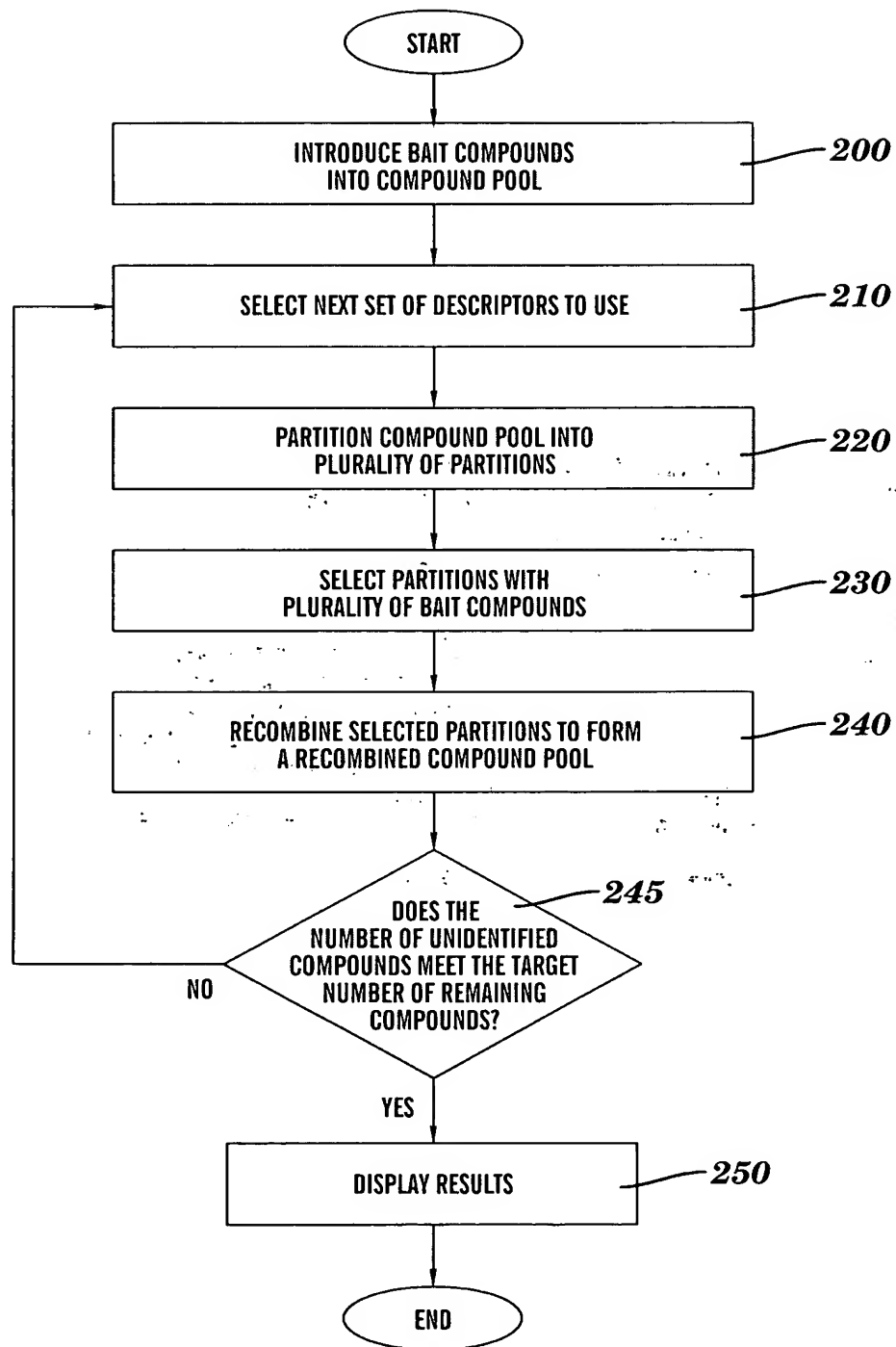


FIG. 12

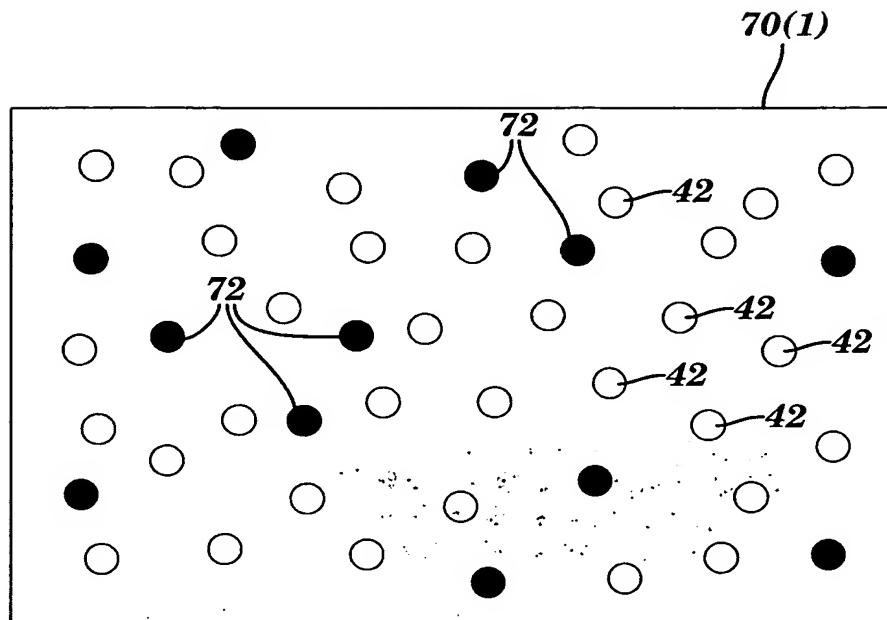


FIG. 13

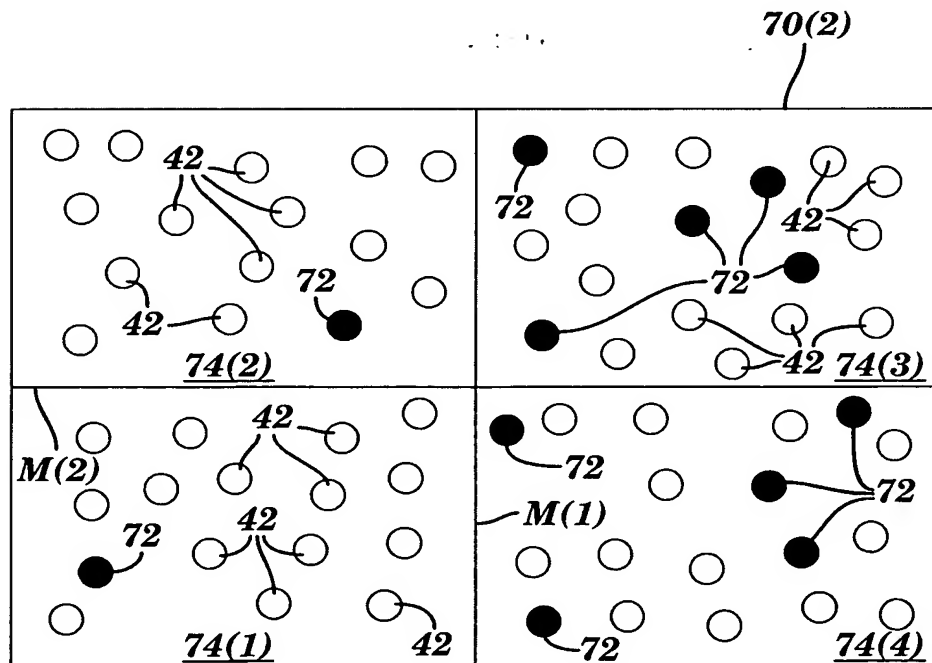


FIG. 14

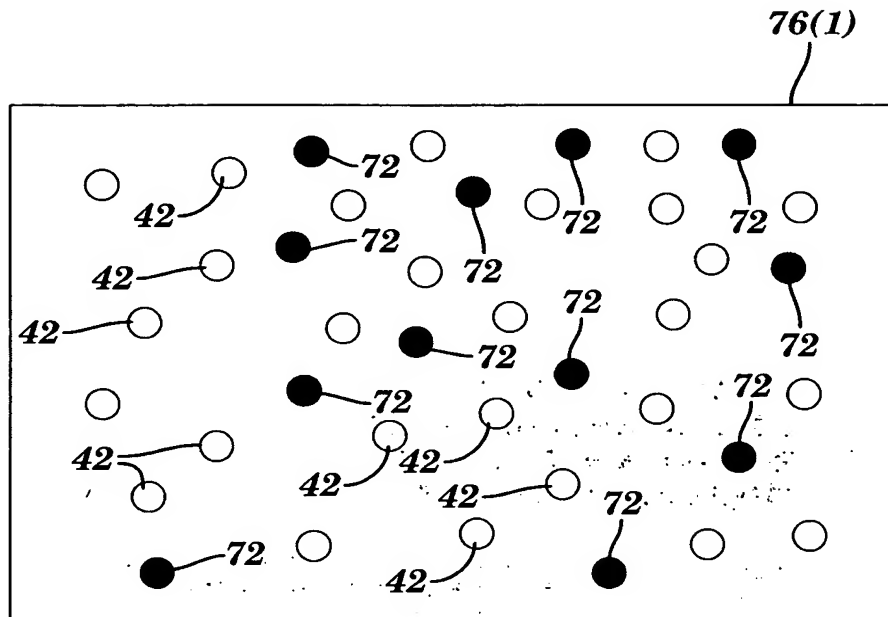


FIG. 15

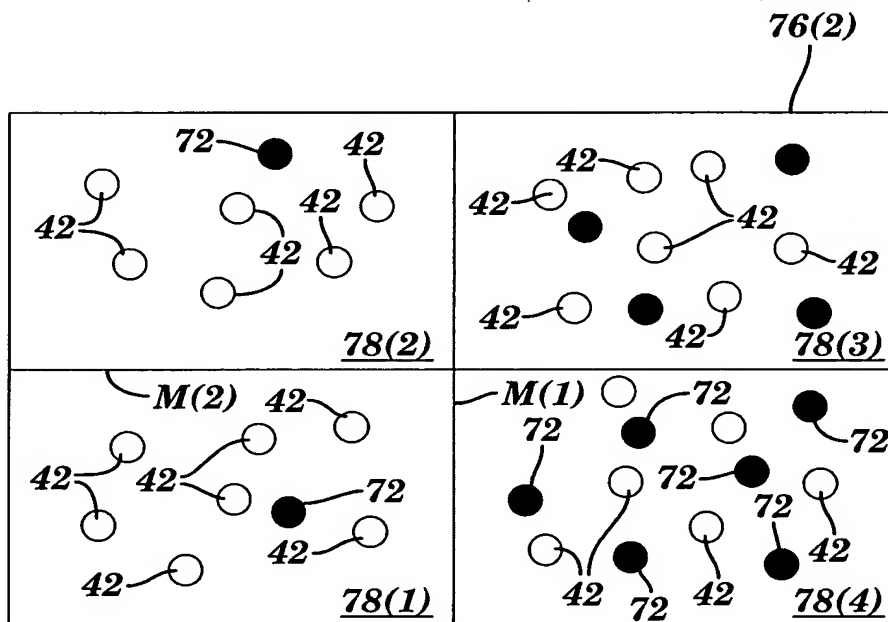


FIG. 16

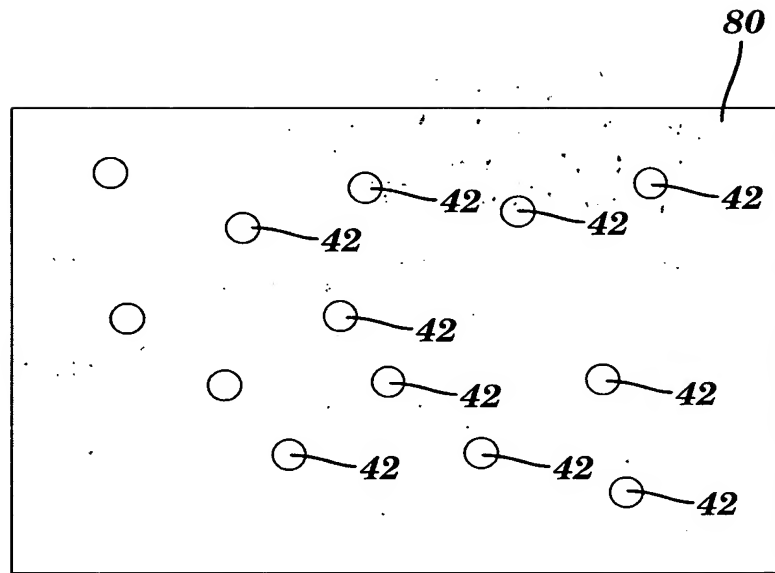


FIG. 17